

SEQUENCE LISTING

<110> Segal, Don
McElroy, Jerry
Chao, Heman
Wong, Wah
Docherty, John
Dickstein, Jodi

<120> Method and Composition for Inhibiting
Cancer Cell Growth

<130> 54800-8023.US00

<140> Not Yet Assigned
<141> Filed Herewith

<150> US 60/397,244
<151> 2002-07-18

<160> 7

<170> FastSEQ for Windows Version 4.0

<210> 1
<211> 35
<212> PRT
<213> Artificial Sequence

<220>
<223> E-coil forming peptide

<400> 1
Glu Val Ser Ala Leu Glu Lys Glu Val Ser Ala Leu Glu Lys Glu Val
1 5 10 15
Ser Ala Leu Glu Lys Glu Val Ser Ala Leu Glu Lys Glu Val Ser Ala
20 25 30
Leu Glu Lys
35

<210> 2
<211> 35
<212> PRT
<213> Artificial Sequence

<220>
<223> K-coil forming peptide

<400> 2
Lys Val Ser Ala Leu Lys Glu Lys Val Ser Ala Leu Lys Glu Lys Val
1 5 10 15
Ser Ala Leu Lys Glu Lys Val Ser Ala Leu Lys Glu Lys Val Ser Ala
20 25 30
Leu Lys Glu
35

<210> 3
<211> 35
<212> PRT
<213> Artificial Sequence

<220>
<223> coil forming peptide

```

<400> 3
Glu Val Glu Ala Leu Gln Lys Glu Val Ser Ala Leu Glu Lys Glu Val
 1           5          10          15
Ser Ala Leu Glu Cys Glu Val Ser Ala Leu Glu Lys Glu Val Glu Ala
 20          25          30
Leu Gln Lys
 35

<210> 4
<211> 35
<212> PRT
<213> Artificial Sequence

<220>
<223> coil forming peptide

<400> 4
Lys Val Glu Ala Leu Lys Lys Lys Val Ser Ala Leu Lys Glu Lys Val
 1           5          10          15
Ser Ala Leu Lys Cys Lys Val Ser Ala Leu Lys Glu Lys Val Glu Ala
 20          25          30
Leu Lys Lys
 35

<210> 5
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> K-coil forming peptide

<400> 5
Lys Val Ser Ala Leu Lys Glu
 1           5

<210> 6
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> E-coil forming peptide

<400> 6
Glu Val Ser Ala Leu Glu Lys
 1           5

<210> 7
<211> 840
<212> PRT
<213> Canavalia ensiformis

<400> 7
Met Lys Leu Ser Pro Arg Glu Val Glu Lys Leu Gly Leu His Asn Ala
 1           5          10          15
Gly Tyr Leu Ala Gln Lys Arg Leu Ala Arg Gly Val Arg Leu Asn Tyr
 20          25          30
Thr Glu Ala Val Ala Leu Ile Ala Ser Gln Ile Met Glu Tyr Ala Arg
 35          40          45
Asp Gly Glu Lys Thr Val Ala Gln Leu Met Cys Leu Gly Gln His Leu
 50          55          60
Leu Gly Arg Arg Gln Val Leu Pro Ala Val Pro His Leu Leu Asn Ala
 65          70          75          80
Val Gln Val Glu Ala Thr Phe Pro Asp Gly Thr Lys Leu Val Thr Val

```

85	90	95
His Asp Pro Ile Ser Arg Glu Asn Gly	Glu Leu Gln Glu Ala Leu Phe	
100	105	110
Gly Ser Leu Leu Pro Val Pro Ser	Leu Asp Lys Phe Ala Glu Thr Lys	
115	120	125
Glu Asp Asn Arg Ile Pro Gly	Glu Ile Leu Cys Glu Asp Glu Cys Leu	
130	135	140
Thr Leu Asn Ile Gly Arg Lys Ala Val	Ile Leu Lys Val Thr Ser Lys	
145	150	155
Gly Asp Arg Pro Ile Gln Val Gly Ser	His Tyr His Phe Ile Glu Val	
165	170	175
Asn Pro Tyr Leu Thr Phe Asp Arg Arg	Lys Ala Tyr Gly Met Arg Leu	
180	185	190
Asn Ile Ala Ala Gly Thr Ala Val Arg	Phe Glu Pro Gly Asp Cys Lys	
195	200	205
Ser Val Thr Leu Val Ser Ile Glu Gly Asn Lys	Val Ile Arg Gly Gly	
210	215	220
Asn Ala Ile Ala Asp Gly Pro Val Asn Glu	Thr Asn Leu Glu Ala Ala	
225	230	235
Met His Ala Val Arg Ser Lys Gly Phe	Gly His Glu Glu Glu Lys Asp	
245	250	255
Ala Ser Glu Gly Phe Thr Lys Glu Asp Pro Asn Cys Pro	Phe Asn Thr	
260	265	270
Phe Ile His Arg Lys Glu Tyr Ala Asn Lys Tyr	Gly Pro Thr Thr Gly	
275	280	285
Asp Lys Ile Arg Leu Gly Asp Thr Asn Leu Leu Ala	Glu Ile Glu Lys	
290	295	300
Asp Tyr Ala Leu Tyr Gly Asp Glu Cys Val	Phe Gly Gly Lys Val	
305	310	315
Ile Arg Asp Gly Met Gly Gln Ser Cys	Gly His Pro Pro Ala Ile Ser	
325	330	335
Leu Asp Thr Val Ile Thr Asn Ala Val Ile Ile Asp Tyr	Thr Gly Ile	
340	345	350
Ile Lys Ala Asp Ile Gly Ile Lys Asp Gly Leu Ile Ala	Ser Ile Gly	
355	360	365
Lys Ala Gly Asn Pro Asp Ile Met Asn Gly Val Phe	Ser Asn Met Ile	
370	375	380
Ile Gly Ala Asn Thr Glu Val Ile Ala Gly Glu Leu Ile Val	Thr	
385	390	395
Ala Gly Ala Ile Asp Cys His Val His Tyr Ile Cys Pro	Gln Leu Val	
405	410	415
Tyr Glu Ala Ile Ser Ser Gly Ile Thr Thr Leu Val Gly	Gly Thr	
420	425	430
Gly Pro Ala Ala Gly Thr Arg Ala Thr Thr Cys Thr	Pro Ser Pro Thr	
435	440	445
Gln Met Arg Leu Met Leu Gln Ser Thr Asp Asp	Leu Pro Leu Asn Phe	
450	455	460
Gly Phe Thr Gly Lys Gly Ser Ser Lys Pro Asp	Glu Leu His Glu	
465	470	475
Ile Ile Lys Ala Gly Ala Met Gly Leu Lys Leu His	Glu Asp Trp Gly	
485	490	495
Ser Thr Pro Ala Ala Ile Asp Asn Cys Leu Thr Ile Ala	Glu His His	
500	505	510
Asp Ile Gln Ile Asn Ile His Thr Asp Thr Leu Asn	Glu Ala Gly Phe	
515	520	525
Val Glu His Ser Ile Ala Ala Phe Lys Gly Arg Thr	Ile His Thr Tyr	
530	535	540
His Ser Glu Gly Ala Gly Gly His Ala Pro Asp Ile	Ile Lys Val	
545	550	555
Cys Gly Ile Lys Asn Val Leu Pro Ser Ser Thr Asn	Pro Thr Arg Pro	
565	570	575
Leu Thr Ser Asn Thr Ile Asp Glu His Leu Asp Met	Leu Met Val Cys	
580	585	590
His His Leu Asp Arg Glu Ile Pro Glu Asp Leu Ala	Phe Ala His Ser	
595	600	605

Arg Ile Arg Lys Lys Thr Ile Ala Ala Glu Asp Val Leu Asn Asp Ile
610 615 620
Gly Ala Ile Ser Ile Ile Ser Ser Asp Ser Gln Ala Met Gly Arg Val
625 630 635 640
Gly Glu Val Ile Ser Arg Thr Trp Gln Thr Ala Asp Lys Met Lys Ala
645 650 655
Gln Thr Gly Pro Leu Lys Cys Asp Ser Ser Asp Asn Asp Asn Phe Arg
660 665 670
Ile Arg Arg Tyr Ile Ala Lys Tyr Thr Ile Asn Pro Ala Ile Ala Asn
675 680 685
Gly Phe Ser Gln Tyr Val Gly Ser Val Glu Val Gly Lys Leu Ala Asp
690 695 700
Leu Val Met Trp Lys Pro Ser Phe Phe Gly Thr Lys Pro Glu Met Val
705 710 715 720
Ile Lys Gly Gly Met Val Ala Trp Ala Asp Ile Gly Asp Pro Asn Ala
725 730 735
Ser Ile Pro Thr Pro Glu Pro Val Lys Met Arg Pro Met Tyr Gly Thr
740 745 750
Leu Gly Lys Ala Gly Gly Ala Leu Ser Ile Ala Phe Val Ser Lys Ala
755 760 765
Ala Leu Asp Gln Arg Val Asn Val Leu Tyr Gly Leu Asn Lys Arg Val
770 775 780
Glu Ala Val Ser Asn Val Arg Lys Leu Thr Lys Leu Asp Met Lys Leu
785 790 795 800
Asn Asp Ala Leu Pro Glu Ile Thr Val Asp Pro Glu Ser Tyr Thr Val
805 810 815
Lys Ala Asp Gly Lys Leu Leu Cys Val Ser Glu Ala Thr Thr Val Pro
820 825 830
Leu Ser Arg Asn Tyr Phe Leu Phe
835 840